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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: Tue Sep 25 09:52:10 EDT 2007

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Application No: 10580901 Version No: 3.0

Input Set:

Output Set:

Started: 2007-09-12 16:33:57.345
Finished: 2007-09-12 16:33:58.037
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 692 ms
Total Warnings: 11
Total Errors: 0
No. of SeqIDs Defined: 11
Actual SeqID Count: 11

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)

SEQUENCE LISTING

<110> Mridula, Sharma
Berry, Carole
Thomas, Mark
Kambadur, Ravi
Bower, Robert Syndecombe

<120> Novel Muscle Growth Regulator

<130> AJPARK39.001APC

<140> 10580901

<141> 2007-09-12

<150> PCT/NZ2004/000308

<151> 2004-11-26

<150> NZ529860

<151> 2003-11-28

<160> 11

<170> PatentIn version 3.1

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<212> DNA

<213> Ovine

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20 25 30

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35 40 45

Leu Gln Thr Gln Thr Pro Pro Pro Thr Leu Gln Gln Pro Ala Pro Pro
50 55 60

Gly Ser Glu Arg Arg Leu Pro Thr Pro Glu Gln Ile Phe Gln Asn Ile
65 70 75 80

Lys Gln Glu Tyr Ser Arg Tyr Gln Arg Trp Arg His Leu Glu Val Val
85 90 95

Leu Asn Gln Ser Glu Ala Cys Thr Ser Glu Ser Gln Pro His Ser Ser
100 105 110

Ala Leu Thr Ala Pro Ser Ser Pro Gly Ser Ser Trp Met Lys Lys Asp
115 120 125

Gln Pro Thr Phe Thr Leu Arg Gln Val Gly Ile Ile Cys Glu Arg Leu
130 135 140

Leu Lys Asp Tyr Glu Asp Lys Ile Arg Glu Glu Tyr Glu Gln Ile Leu
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Asn Thr Lys Leu Ala Glu Gln Tyr Glu Ser Phe Val Lys Phe Thr His
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180 185 190

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cccgccccgc ccggcagcga ccggcgctt ccaactccgg agcaaat ttcagaacata 240

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Thr	Leu	Thr	Ala	Pro	Ser	Ser	Pro	Gly	Ser	Ser	Trp	Met	Lys	Lys	Asp	115	120	125	
Gln	Pro	Thr	Phe	Thr	Leu	Arg	Gln	Val	Gly	Ile	Ile	Cys	Glu	Arg	Leu	130	135	140	
Leu	Lys	Asp	Tyr	Glu	Asp	Lys	Ile	Arg	Glu	Glu	Tyr	Glu	Gln	Ile	Leu	145	150	155	160
Asn	Thr	Lys	Leu	Ala	Glu	Gln	Tyr	Glu	Ser	Phe	Val	Lys	Phe	Thr	His	165	170	175	
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